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# Overview of Congenital, Senile and Metabolic Cataract

**Keywords:** Cataractous lens, Congenital, Senile cataracts, Genetic factors, Metabolic pathways

#### Abstract

Opacity of the lens is a major public health problem of unknown causes and impaired vision of a large number of peoples. There are different forms of cataractous lenses including congenital, senile and metabolic associated ones. Although the congenital and senile cataractous lenses were markedly different in its origin, developmental aspects and genes involved, the patho-physiological pathways seemed to be similar. The present review summarized the causes of cataractous lenses such as genetic and abnormal conformational changes of glycation end products. The alterations of proteins, lipids, minerals and antioxidants defense were illustrated and discussed.

# Introduction

The lens is avascular biconvex ellipsoid transparent tissue, present in front of the anterior chamber of the eye; just behind the cornea. It is an ectodermal organ, developed at the 25<sup>th</sup> day of gestation. By the second month of intra-uterine growth, the spherical criteria structure of the lens is developed with characteristic anterior and posterior pole and dorsally outlined by lens epithelium. The lens fibers developed and matured after birth with subsequent formation of crystalline protein especially in adult state [1].

The cornea is affected by continues light exposure which may be reflected in the structure of the lens with age. The lens comprises three main regions; capsule, lens epithelium, and lens fibers. The capsule is made up of dense connective tissue. The lens epithelium; lined the surface and it comprises a simple cuboidal epithelium. It possesses a vital function by regulating the homeostatic functions allowing the permeability of ions, nutrients, and osmolarity to the aqueous humor. It develops earlier during utero life in the 5th-6th week of gestation and maintains its normal integrity throughout [2]. The entire body of the lens is composed of concentric layers of tightly backed lens fibers interdigitated with each other by a ball and socket. Glucose represents the primary energy source of the lens tissues [3]. Sodium/potassium adenosine triphosphatase and calcium adenosine triphosphatase promote the osmolarity of the lens fibers [4].

Damage of the lens fibres lead to the formation of clouding in the lens and scattering of light, the predictor of cataract. It increases with age with a prevalence of about 50% in over 80 years old individuals [5]. There are different forms of cataracts; namely congenital [6] agerelated cataract [7] and metabolic cataract [8].

# **Congenital & Senile Cataract**

Cataract is a clouding of the eye's natural lens which causes visual impairment. More than 17 million people are blind due to cataract and 2800 new cases are known throughout the world daily [5].

Congenital cataract are developed during intra-uterine growth as a result of environmental and genetic factors and observed after parturition. Approximately half of all congenital cataracts are

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**Review Article** 

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idiopathic [9] and particularly, a cause for unilateral cataract is rarely found [10].

Congenital cataracts are of different genetic type, and more than 25 loci and genes on different chromosomes are involved in the disease [11]. Rahi and Dezateux reported a similar genotype in 27% of children with bilateral congenital cataract [12].

In age related cataract, many changes have been found to have occurred in the lens structure which facilitated the opacities of lens fibres. The epithelia of nuclear, posterior subcapsular, mature, mixed, hypermature, and black cataracts of male and female patients revealed that the 56% superimposed epithelial cells are probably the source of increased and altered cell activity. Senile cataract is classified into; mature or immature, nuclear cortical and posterior subcapsular cataract [13]. Investigation of 233 cataractous patients, revealed apparent increased incidence of nuclear, cortical, and posterior subcapsular cataract in older patients. These were correlated with lower expression of SIRT1 in patients [14].

# **Mechanism of Cataract Formation**

Metabolic diseases were found to be associated with the activation of the sorbitol pathway [15] that enhanced the progress of cataract formation. Liberation of free radicals is more common as a result of these diseases. *In vivo* and *in vitro* studies revealed that hydrogen peroxide ( $H_2O_2$ ) is related to the accumulation of A $\beta$  in the lenses of rats and causes lens opacification [16]. The incidence of cataractous lenses was markedly increased among patients with metabolic diseases such as hypertension [15] and diabetes [17]. Hypertensive patients showed significantly higher nitrite levels in their cataractous lenses were detected in 26 (44.1%) especially that of posterior subcapsular cataracts [18,19].

Moncaster et al. reported increased amyloid accumulation and supranuclear cataracts in the ocular lenses of patients with Alzheimer's disease [20].

There are different pathways for the formation of cataract (Figure 1), they include:

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Figure 1: Chart illustrates the mechanism of cataract formation. Abbreviations: CE: Corneal Epithelium; CO: Ciliary Organ; E: Endothelium; LC: Lens Capsule; LE: Lens Epithelium; LF: Lens Fiber; LN: Lens Nucleus; St: Stroma.

# Polyol pathway

In diabetic cataractous lens, there was a marked increase of damaging lens epithelium [21] associated with over production of free radicals and decreased capacity of antioxidant defense system. These were associated with a depletion of the activities of aldose reductase and sorbitol dehydrogenase while G6PD and glutathione system enzyme activities were found to be lower in cataractous lenses from diabetes [22]. Aldose Reductase is a key enzyme involved the reduction of glucose into sugar alcohol sorbitol, which is metabolized to fructose by sorbitol dehydrogenase. Sorbitol was found to accumulate in lens cells causing osmolysis, leakage of glutathione, myo-inositol, the generation of free radicals which contributed to diabetic complication such as cataract [23,24]. The enzyme is more active in diabetes and implicated in cataractous lenses [25,26].

Sugar cataract formation was found to have resulted from lenticular sorbitol accumulation. In the lens, increase of aldose reductase is contradicted with a decrease of the activity of sorbitol dehydrogenase [27]; it increases the accumulation of sorbitol rapidly than it is converted to fructose [28] which induces apoptosis of lenticular epithelial cells and subsequent cataracts [29]. Diabetes was found to express poly(ADP-ribose) polymerase(PARP) in lens [30] which in turn led to NAD<sup>+</sup> depletion and energy failure and also contributed to necrosis and apoptosis [31].

Intracellular accumulation of sorbitol is associated with marked increase of oxidative stress in the endoplasmic reticulum (ER), the principal site of protein synthesis, it initiates the liberation of free radicals and also involved in the breakdown of lens fibers [32,33]. Over-expression of aldose reductase also enhanced the expression of both extracellular signal-regulated kinase (ERK1/2) and c-Jun N-terminal (JNK1/2), which are involved in apoptosis [34].

In sugar cataracts, the initial swelling brought about by polyol accumulation leads to an imbalance in the pump-leak equilibrium, increasing sodium and chlorine content. Genetic mouse cataract was found to exhibit the imbalance of the pump-leak system which appeared to be initiated by a deficiency of Na-K ATPase which facilitates sodium retention and osmotic swelling [35].

Nagai et al. reported a decrease in the activity of  $Ca^{2+}$ -ATPase in the lenses of Shumiya cataract rat (SCR) and Ihara cataract rat (ICR), which is concurrent with the development of cataract [36]. The expression of cytochrome c oxidase (CCO)-1 mRNA and CCO activity in UPLR lenses was found to have decreased during cataract development. Both nitric oxide (NO) and lipid peroxide were markedly increased in the lenses of UPLR, SCR and ICR with opaque lenses. Increased liberation of NO may enhance lipid peroxidation resulting in the inhibition of  $Ca^{(2+)}$ -ATPase and elevation in lens  $Ca^{(2+)}$ , leading to lens opacification in ICR/f rats.

**Glycation end product (AGE)**: AGE is a non-enzymatic glycation of lens proteins by reducing sugars (glucose, galactose etc) resulting in the formation of protein aggregates that precipitate in the lens [37,38] and represent a causative factor of damaging lens epithelial cells (LECs), the promoter of cataract formation [39]. The receptor for advanced glycation end-products (RAGE) and proliferating cell nuclear agents (PCNA) are over-expressed in old-age and diabetes which is involved in cataract formation [40,41]. Advanced glycation end products (AGEs) including carboxymethyl lysine (CML) was found to be increased in diabetic and senile cataractous lenses [42].

The urea-insoluble fraction, showed the highest levels of AGEs [43]. Pentosidine, CML, and imidazolone were markedly increased in cataractous lenses [44]. *In vitro* and *in vivo* studies revealed AGE induced apparent apoptosis of LECs through enhancement of accumulation of argpyrimidine and nuclear factor- kappaB (NF- $\kappa$ B). The ratio of Bax to Bcl-2 protein levels was also increased [39].

Cataractous lenses were markedly higher in women especially those with diabetes [45]. In cataractous lens of 21-week-old male ZDF rats, increased apoptosis of LECs was correlated with upregulated expression of iNOS mRNA and protein, causing the accumulation of glycated proteins in their cytoplasm, meanwhile activated NFkappaB was mainly detected within their nuclei [46].

# **Genetic Role of Cataract Formation**

There are multifactorial agents that contribute in the development of cataractous lenses as well as between congenital and age-related cataracts. Congenital cataract affects childhood, and different intrauterine factors are involved (Figure 2). The genetic incidence accounts for 8.3 and 25% cataract. Congenital cataracts may result from chromosomal abnormalities such as neurofibromatosis type 2 or homeobox gene PITX3 [47,48] classified congenital abnormalities into mature, polar (anterior or posterior), zonular (nuclear, or lamellar), and capsular or membranous.

Zhu et al. attributed the congenital cataract to the mutation

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Figure 2: Diagrammatic representation the incidence and genetic role in congenital and senile cataract formation.

Abbreviations: CE: Corneal Epithelium; CO: Ciliary Organ; E: Endothelium; LC: Lens Capsule; LE: Lens Epithelium; LF: Lens Fiber; LN: Lens Nucleus; St: Stroma.

of connexin 50 of a valine residue with alanine at codon 44 [49]. In cataract lenses of KO-Dp71 mice and wild-type (wt), Fort et al. reported increased incidence of cataracts with subsequent lack of Dp71 (dystrophin gene of Duchenne muscular dystrophy) [50].

Khan et al. observed posterior cortical lenticular opacities with over-expression of a novel missense mutation in KCNJ13 in 12-yearold Saudi Arabian girl with nystagmus [51].

CF1/b mouse strain (an outbred colony) reared at Osaka Prefecture University was found to develop cataracts at 14 d old in the form of vacuolated lens epithelial cells, swollen lens fibers, pyknotic nuclei, and vacuolation of the lens cortex. Cataractous lens was found to result from autosomal recessive mutation genes in mouse chromosome 16 between D16Mit5 and D16Mit92 and between D16Mit92 and D16Mit201 [52].

Lens syntaxin 3 gene interfered with lens protein structure and function. Mutation of this gene results in the development of autosomal recessive congenital cataract [53]. Inherited forms of cataract are characterized by mutation in the gene of cation channel and member-3 (TRPM3, melastatin-2). Genome-wide linkage analysis mapped the ocular disease locus to the pericentric region of human chromosome 9 [54].

Although congenital cataracts may alter vision; sparse individuals

contributed to the pathogenesis of age-related cataract (ARC) [60].

Oxidative stress represents the main factor responsible for lenticular cataract. Glutathione-S-transferase (GST) catalyses the nucleophilic addition of the thiol of GSH to electrophilic acceptors, it is important for protecting tissues from oxidative damage. In humans, GSTT1 and GSTM1 deletion genotypes are associated with a variety of ophthalmic diseases. The frequency of GSTM1 affected individuals was markedly increased in MT cataracts followed by NC, CC and PSC types [61].

125b and p53 expression [59]. Oxidative stress and DNA damage

are affected, age-related cataracts are responsible for blinding 17

occurred in the structure of the lens which facilitated the opacities

of lens fibres. Telomerase represents one of the genetic factors. It is a naturally occurring enzyme that maintains telomeres and prevents

them from shortening during cell division. Telomerase consists of at

least two components: an RNA template (hTR), which binds to the telomere and a catalytic subunit (hTERT) with reverse transcriptase activity, which adds a specific DNA sequence to the chromosome ends.

During the formation of cataract, telomeres attained a considerable atrophy in lens epithelial cells due to increased oxidative stress of the lens cell membranes and biomolecules [56,57]. Cataractous patients possessed the presence of a large number of micronuclei (MN) in

MicroRNA-125b (miR-125b) is responsible for lens epithelial cell apoptosis. *In vitro* studies of miR-125b revealed increased incidence of human lens epithelial cell apoptosis through assessments of p53. In age-related cataract, there was an inverse relationship between miR-

their epithelia, manifesting epithelial damage [58].

In age related cataract, many changes had been found to have

million people worldwide [55].

Peroxiredoxin 6 (Prdx6) is a new family of antioxidants which regulates gene expression and function by scavenging reactive oxygen species and protects lens epithelial cells of rat. Oxidative stress was found to have contributed in the formation of nuclear or cortical cataract through a decrease of Prdx6 in cataractous lenses especially during aging [62].

Nuclear transcriptional factor, NF-E2-related factor 2 (Nrf2), plays an integral role against oxidative stresses by promoting 20 different antioxidative enzymes. Kelch-like ECH associated protein 1 (Keap1) targets and binds to Nrf2 for proteosomal degradation. Palsamy et al. detected significant levels of demethylated DNA in the Keap1 promoter in the diabetic cataractous lenses [63]. Human lens epithelial cell lines (HLECs) treated with a demethylation agent, 5-aza-2'deoxycytidine (5-Aza), showed 10-fold higher levels of Keap1 mRNA, 3-fold increased levels of Keap1 protein, which enhanced the liberation of free radicals and increased cell death. Decreased Nrf2 activity depleted the transcription of many antioxidant enzyme genes and altered the redox-balance towards lens oxidation. During agingrelated cataract, there was a considerable reduction of protein and gene expression level of nuclear transcriptional factor, NF-E2-related factor 2 (Nrf2) in lenses. Over-expressed Keap1 protein is responsible for decreasing Nrf2 by proteasomal degradation, thereby suppressing Nrf2-dependent stress protection. Therefore, decreased activity of Nrf2 restrained the transcription of its downstream antioxidant enzyme and led to the failure of the antioxidant system; ultimately

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# leading to the formation of ARCs [64,65].

In experimental animals, different pathways are involved. Transgenic mice were found to express ectopic Col4a3 and Col4a4 genes in the lens which have the capability of activating IRE1, ATF6, and PERK associated with the expansion of the endoplasmic reticulum and attenuated protein translation. These were associated with damaging lens epithelium and accumulate terminally unfolded proteins [66]. Following the study of three mice lens epithelial cells (LECs) 17EM15, 21EM15 and  $\alpha$ TN4, Terrell et al. reported that the expressed genes Foxe3, Pax6, Anxa4 and Mcm4 were up-regulated in lens epithelial cell lines (LEC), compared to lens fiber cells [67]. All the three LECs exhibited down-regulation of fiber cell-expressed genes Crybb1, Mip and Prox1 as well as over-expressed cataract-associated genes, such as Dkk3, Epha2, Hsf4, Jag1, Mab2111, Meis1, Pknox1, Pou2f1, Sfrp1, Sparc, Tdrd7 and Trpm3.

# Lipid and Cataracts

Lipids are of functional importance to lens. About 172 lipid species were identified mainly as sphingomyelins, phosphatidylcholines, and phosphatidylethanolamines. Also, 20 phosphatidylcholines, 6 phosphatidylethanolamines, and 4 phosphatidic acids were also detected [68]. The lens transmit light through thousands of cellular membranes rich in phospholipids especially dihydrosphingomyelin. Most of the lipids are associated with proteins. Sphingolipids represent 60% of human lens and represents the main integral part in its transparency. Lens glycolipids are composed of 4-sphingenine (sphingosine) to which carbohydrates are bound [69] and represents 1% of the total human lens lipid but are of valuable importance for differentiation of lens epithelial cells to lens fibers [70].

Lipids represent the main structural components of the lens membranes and have profound effect on its fluidity. Cataractous lens involves protein aggregation as a result of losing stability in protein conformation. Age-related changes in lipid composition could be a contributing factor for altered protein-lipid interaction leading to protein aggregation and cataract formation [71]. The lens membranes exhibited three distinct lipids such as the bulk, nuclear, and trapped lipid domain. The cholesterol bilayer was detected in the cortical and nuclear lens lipid membranes, and missing in the intact membranes. The amount of lipids combine proteins was increased in the nuclear than in cortical membranes facilitating rigidity of nuclear membranes and increasing permeability in the cortical ones [72].

Cataractous lenses were found to exhibit membrane derangement, widespread of vacuoles and clusters of highly undulating membranes. Damage of the lens fiber cell membranes impairs vision and interferes with light-scattering which causes lens opacity. Phospholipid molecules modified by oxygen accumulate in the lipid bilayer, they have been found to change their geometry and impair lipid-lipid and protein-lipid interactions in the lenticular fiber membranes [73]. Human cataractous lenses contain the long chain base glycolipids (C18a sphingosine (sphinganine). The major fatty acids were C160, C24:1 and C24:0, and monounsaturated fatty acids accounted for 40-55% the total fatty acids [74]. Siddique et al. reported alterations of phospholipid and protein moieties in lens fibers, disrupting function and leading to cataract formation (Figure 3) [75].

The content of palmitic acid showed apparent increase in



Figure 3: Diagrammatic representation the role of lipids in senile cataract formation.

Abbreviations: CE: Corneal Epithelium; CO: Ciliary Organ; E: Endothelium; LC: Lens Capsule; LE: Lens Epithelium; LF: Lens Fiber; LN: Lens Nucleus; St: Stroma.

cataractous lens when compared with marked depletion of oleic acid, linolenic acid and docosahexaenoic acid [76].

Huang et al. detected increased amount of sphingolipids (dihydrosphingomyelin and sphingomyelin) compared with apparent depletion of glycerolipids (phosphatidylcholine and two phosphatidylethanolamine) in cataractous lens [77].

Lipid peroxidation (LPO) is a pathogenic factor in cataract. It includes diene conjugates, lipid hydroperoxides, oxy-derivatives of phospholipid fatty acids and present in the lipid moieties of aqueous humour and lenses of senile patients [73]. Plasma membranes of the ocular lens are rich in fiber junctions which in turn have 20-40% of total lipid content compared with that of the total plasma membrane. In the bovine nucleus, the cholesterol/phospholipid molar ratio was markedly increased in the fiber junctional-enriched membrane than in the total plasma membrane, suggesting a special association of cholesterol with bovine nuclear fiber junctions [78]. In cataractous human lenses, the ratio of cholesterol to phospholipid (Chol/PL) attained a considerable increase in lens membranes.

Exposure of membranes isolated from transparent human lenses to the free radical generator were found to produce 7 alphahydroxycholesterol (6%), 7 beta-hydroxycholesterol (19%), 5 alpha, 6 alpha-epoxycholestanol (1%) and 7-ketocholesterol (74%) as

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major oxidation products. Cataractous lenses possessed moderate amounts of 7 beta-hydroxycholesterol, 7-ketocholesterol, 5 alpha, 6 alpha-epoxycholestanol, 20 alpha-hydroxycholesterol and 25-hydroxycholesterol compared with no-detectable amounts in clear lenses [79]. Studies have shown that cholesterol contributes to the impermeability of oxygen across the lens membranes [80]. Maintaining low levels of oxygen is important to the lens to prevent oxidation and cataract formation. One of the main function of the mitochondria, present in the epithelium and outer cortical fibers [81] is to degrade oxygen [82]. The cholesterol-related impermeability of the membranes to oxygen may help to keep oxygen in the outer regions of the lens long enough for the mitochondria to degrade it.

The etiology of lipids in cataract formation led Deeley et al. to investigate the variation of eye lipid contents in humans and experimental animals [83]. The authors found that the most abundant phospholipids in all the lenses examined were cholinecontaining phospholipids. Human lens showed wide variations of lens lipid components compared with rat, mouse, sheep, cow, pig and chicken. Sixty six percent (66%) of the total phospholipid in Homo sapiens was sphingomyelin (dihydrosphingomyelins). The abundant glycerophospholipids in human lenses were phosphatidylethanolamines and phosphatidylserines. Experimental and domestic animals showed abundant phosphatidylcholines in their lenses. Several neutral and acidic glycosphingolipids were observed in rat lens epithelia, cortex and nucleus. Immunohistochemical staining revealed apparent concentration of gangliosides GM3 and GM1 in the anterior lens epithelial cells and the cortex, with gradual decrease toward the lens nucleus. GD3 appeared more intense in the lens nucleus than in epithelial cells [84]. Adult rat lens were found to have a complex ganglioside consisting of six major components identified as GM3, GD3, GD1a, GD1b, GT1b, and GQ1b based on their reactivity to anti-GM1. The relative concentration of GT3 in total gangliosides of the eye lens was highest among neural and extra-neural tissues examined. Administration of streptozotocin to rats caused a severe reduction in the GD3 content in the eye lenses within three days prior to other gangliosides alterations [85]. Emory mouse cataracts undergo protein oxidation by the formation of hydrazone with 2, 4-dinitrophenyl hydrazine. The lipid oxidation was observed from a depletion of oleic acid and over-expression of ketoacids [86]. The relative and absolute amount of sphingolipids, including dihydrosphingomyelin and sphingomyelin, increased with age parallel with the development of cataractous lenses. Increased membrane stiffness led to light-scattering, reduced calcium pump activity, altered protein-lipid interactions, and decreased fiber cell elongation [77].

# **Protein and Cataracts**

The lens protein attained nearly 33% of its net weight. Crystallins are the main protein components in the lens reaching up to 90% of total soluble protein. Other proteins form the cytoskeletal and membrane structure such as actin, filensin and spectrin, transporters and channel proteins and junctional proteins. There are three distinct forms of crystallins;  $\alpha$ -,  $\beta$ - and  $\gamma$ -crystallins.  $\alpha$ - and  $\beta$ -crystallins are present in oligomers, compared with the monomer form of  $\gamma$ -crystallin [87]. In adults, the nucleus represents the major part of the lens. The nucleus is concerned with cell communication and many enzymes involved in

metabolism, protein synthesis and degradation [88].

In old age, there is a marked disruption of lens disulphide protein aggregates and conformation which is unique to cataract. In cataracts, 60% of the methionine is bounded to the lens membrane. Most of the disulfide amino acids bind with glutathione to form  $\alpha$ -crystallin via oxidative changes [89]. Homozygous alpha alanine-cysteine mutant mice, possessed apparent characteristic loss of cytological structure of lens epithelial and fiber cells such as actin filaments and mitochondria (alpha, beta, and gamma classes) [90]. Alpha-A crystallin represents the major protein of the lens and is composed of a mixture of cysteine sulfhydryl and half-cysteine disulfide groups. Oxidation of cysteine sulfhydryl groups may be involved in opacification of lens [91].

Crystallins are categorized in three forms  $\alpha$ ,  $\beta$ , and  $\gamma$ . The  $\beta$ -crystallins remain the most elusive form due to their greater number of subunits and possible oligomer formations and changes during aging. Deamidation of  $\beta$ -crystallins occurred during aging in cataractous lenses and  $\alpha$ -crystallin chaperone impaired the cataractous changes [92]. The cysteine is characterized by having disulfide bond with the low molecular weight thiol in the lens. The appearance of gamma-Glu-Cys may be coincident with biochemical abnormalities preceding cataract formation. This protein modification may result from changes in the GSH biosynthetic pathway within the lens [93,94]. In cataractous patients with or without diabetes, there were marked aggregations of two types, soluble and insoluble proteins. Higher levels of m-Tyr, o-Tyr, DOPA and Phe/protein ratio were



Figure 4: Diagrammatic representation the role of protein in senile cataract formation.

Abbreviations: CE: Corneal Epithelium; CO: Ciliary Organ; E: Endothelium; LC: Lens Capsule; LE: Lens Epithelium; LF: Lens Fiber; LN: Lens Nucleus; St: Stroma.

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markedly detected in cataract formation (Figure 4) [95].

Opacification of the ICR/f mutant lenses was found to be accompanied by changes in crystallin structure and composition, including several deletions of the N-terminals of beta-crystallins and low molecular weight alpha-crystallins [96].

Cataractous lens proteins exhibited apparent decrease in gamma-, betaB1-, betaA3-, and betaA4-crystallin content, accompanied with some increase in alpha-crystallin (or its aggregate). Higher molecular weight proteins were also observed in the form of cross-linked dimers (43 to 55 kDa) of beta-crystallins [97]. Deletions of the N-terminals of beta-crystallins and low molecular weight alpha-crystallins were detected in cataractous ICR/f mutant lenses [96]. Aggregation of chromophores and insoluble crystallin led to the formation of a brown nuclear cataract while cortical type resulted from impairing of membrane permeability and altered enzyme function [88].

Linetsky et al. stated that ascorbic acid, but not glucose, fructose, ribose or erythrulose, facilitated the aggregation and glycation of calf lens crystallins [98]. Dipeptidyl peptidase III was markedly increased in cataractous lens to an apparent 45.5-fold over that of the original aqueous extract [99]. There are two forms of cataractous lenses, one with a low internal sodium and calcium content which behaves in a manner similar to normal lenses. Elevated level of sodium ion was found to be involved in alterations of amino acid accumulation, however calcium ions seem to play a critical role in the disturbance of lens protein synthesis [100]. Water-insoluble, high molecular weight protein represents the major component of human nuclear cataracts [101].

AGEs formation resulted from oxidation reaction between reducing sugars and protein which caused lens opacification. Diabetes showed more glycation in patients suffering from diabetes compared with non-diabetic. Pyrraline is an advanced Maillard reaction formed by oxidation of glucose with lysine residues on proteins. This reaction involves an intermediate metabolite, 3-deoxyglucosone which is markedly increased in diabetic lenses [38,102].

Lens proteins possessed the higher levels of dideoxyosones (DDOs): intermediates in the synthesis of advanced glycation endproducts (AGEs), such as pentosidine and glucosepane [103]. Deamidation of glutamine and asparagine residues was observed in older lenses [104].

One of the possible roles of inducing lenticular diabetic cataract is the capability of glycated proteins producing reactive oxygen species (ROS), which oxidized tryptophan (Trp) into kynurenines [105,106]. There were marked differences in the amounts of oxindolealanine (OIA, tryptophan oxidation product) in the nucleus versus the cortex in human cataractous lenses [107]. Lenticular levels of Indoleamine 2, 3-dioxygenase (IDO) activity, IDO mRNA, IFN-gamma mRNA, oxidize tryptophan (Trp) and kynurenic acid (KYNA) increased significantly in 60 days diabetic rats [106]. Histidinoalanine and lanthionine were the most abundant dehydroalanine crosslinks in both water-soluble and -insoluble lens proteins in cataractous lenses of Indian patients [108].

Another view of cataract formation is believed to have resulted from a decrease in the antioxidant defense of the lens, particularly the vitamin C concentration. Upon oxidation, vitamin C was found to have combined with glucose to form protein glycation. The lens vitamin C concentration significantly decreased with cataract severity, especially in severe brown cataracts. The peptide tryptophan content was stable but the tryptophan to tyrosine ratio decreased and was highly correlated to the ascorbic acid concentration [109]. Taurine and total protein were detected in cataractous lenses and involved in altering the structural integrity and permeability of lens membrane to protein and amino acids [110]. Taurine showed antioxidant capacity and was markedly depleted in diabetic cataract. The culture of lenses in high glucose medium increased the weight and opacity of the lenses due to increase of carbonylated protein level, and decrease of glutathione (GSH) content [111].

# **Minerals and Cataract**

# Sodium & potassium

Electrolyte balance is of vital importance to the lens transparency in mammals including humans. Its permeability depends on different ion channels such as potassium, sodium, chloride, and calcium channels. The ions may move across a number of electrical silent system such as Na<sup>+</sup> /H<sup>+</sup>, Na<sup>+</sup> /Ca<sup>2+</sup> HCO<sup>3-</sup>/Cl- exchange mechanisms as well as active transporters such as Na-K-ATPase [112,113]. Na, K-ATPase activity plays an important role in maintaining the functional activity in lens. Two apparent roles of the enzyme activity were observed more in the epithelium than in the lens fibers. During aging, oxidation and glycation of lens fibers may decrease Na, K-ATPase activity which is markedly detected in the nuclear region [114,115]. There is a close association between abnormal elevation of lens sodium and the opacification of the human lens cortex [116]. Aging has also led to alteration of Na/K permeability toward the center of the lens as well as impairing the functional activity of the Na-K-ATPase to hydrolyze adenosine triphosphate [117]. Increasing extracellular K<sup>+</sup> was found to depolarize the lens potential, reducing and reversing the magnitude of the net current densities around the lens [118].

There are two forms of cataractous lenses, one with a low internal sodium and calcium content behaving in a similar manner to normal lenses, and the other with high sodium and calcium contents which showed a markedly reduced ability to accumulate amino acid and synthesized low molecular weight protein. Sodium ion was found to be involved in alterations of amino acid accumulation, however calcium ions seems to play a critical role in the disturbance of lens protein synthesis and also protein-protein interaction [119].

# Magnesium

Magnesium represents one of the main elements in regulating the lens function. It maintains the function of more than 350 enzymes by regulating the intracellular ionic environment of the body ATPase. Its deficiency led to impairments of ATPase functions leading to marked increased of intracellular calcium and sodium and decrease intracellular potassium concentration and development of cataract. Consequently, magnesium deficiency has also contributed to the increase of oxidative stress and inducible NOS stimulation that can initiate the progress of cataract, glaucoma and diabetic retinopathy [120,121].

# Zinc

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Zinc plays an integral role in maintaining normal ocular function. The zinc content of human lenses was significantly elevated in mature cataracts compared to cortico-nuclear cataracts. The increase of the lens zinc content was markedly increased with the change in lens coloration from light to dark brown colouration. Diabetic patients were found to show both increased zinc and iron contents in the lens. Zinc and iron were markedly increased in cataractous lenses [122].

Lens opacification in diabetic patient was found to possess increased level of Cu content than Zn and Fe content [123]. Patients with either pseudoexfoliative or senile cataract showed increased serum level of both iron and copper [124].

 $\alpha$ -Crystallin, is a member of the small heat shock protein family present in the mammalian lens. The stability of  $\alpha$ -crystallin is increased in the presence of Zn(<sup>2+</sup>). H79, H107 and H115 of  $\alpha$ A-crystallin and H104, H111 and H119 of  $\alpha$ B-crystallin are identified as the Zn(<sup>2+</sup>) binding residues [125]. Development of lens opacities was associated with lack of copper-zinc superoxide dismutase which participated in an antioxidant capacity [126].

There is a close association between mineral and antioxidant defense. Cataractous lens possessed a decrease of Cu/Zn- and Mn-SOD activity and Cu/Zn-SOD transcript. The decrease of superoxide dismutase activity in cataractous lenses was associated with the decreased level of mRNA transcripts and their protein expression [127].

# Iron

Iron (Fe) is an essential trace element for ocular homeostasis. It was found to play a great role in regulating L-cystine uptake and consequently promoting the activity of aconitase in production of glutathione in lens epithelium and retinal pigmented epithelium that led to elevation of the antioxidative capacity against oxidative stress [128]. During the development of lens epithelial cells into lens fiber, both ferritin H and L chains were altered with increase in the mass of lens fiber [129]. Iron pathway may occurred through different mechanisms; (i) iron is actively transmitted from the retina to the aqueous humor by a ferroportin/ferroxidase-mediated process through Müller cells, (ii) binding of iron with vitreal transferrin and then diffusing towards the lens, (iii) Incorporation of iron/ transferrin complex with lens extracellular space and infiltrated into the epithelial-fiber interface, (iv) iron/transferrin complex is endocytosed by epithelial cells and drained with the aqueous humor into systemic blood circulation for recycling [130].

Transferrin and Fe concentrations were found to be increased in the intraocular fluids during inflammation and pathological conditions. There was a close relationship between the amount of Fe-transferrin and development of cataractous lenses [131]. The ferritin levels in the cataractous lens nuclei appeared to be markedly increased than that of the cortex due to the presence of ferritin within an insoluble protein fraction of the homogenized lenses. Both nuclei and cortices showed heavily redox-active metals staining [132]. Hyperferritinemia cataract formation was found to result from a mutation in the L-ferritin gene (C33T) which interferes with function of the L-ferritin transcript in an RNA gel shift assay [133].

Cu and Fe lacking a non protein-binding condition, initiate the liberation of reactive oxygen species, thereby affecting tissue and

cell structures [134]. Abnormal metabolism of Zn, Cu and Fe may be influenced in diabetic complications [134], especially Cu which reached a high peak in the blood of diabetic patients [135] compared with low peak of zinc [136,137] and iron [138].

Iron was found to have contributed in ocular diseases, including glaucoma, cataract, AMD, and intraocular hemorrhage [139]. Numerous genes encoding proteins involved in iron transport and homeostasis are promoted by hypoxia inducible factor. Low oxygen level was found to stimulate the lens epithelial cells (LEC) to trap the cytosolic iron and increased the risk of iron inducing the formation of reactive oxygen species (ROS) and oxidative cell damage [140].

Diabetic cataractous lenses of rats were found to possess increased level of Zn and Fe and decreased level of K [141]. In senile cataractous lenses, Fe, Al, Zn and Ca levels were markedly increased. The calcium/ phosphorous ratio was 50 times (1:0.02) greater than in clear lenses [142]. Senile cataractous lenses were found to possess elevated level of nickel and iron with concomitant depletion of chromium, manganese and aluminum, the promoter of cataract formation [143]. Li et al. detected degenerated lens fibers in nuclear cataract associated with aggregation of gamma-crystallin in lens fibers lacking actin filaments and increased calcium concentration of homozygous mutant mouse [144].

# Calcium

Calcium plays a great role in large numbers of cellular regulatory pathways. The transmembrane influx of  $Ca(^{2+})$  through Ca(v) channels has contributed to the biological function of human lens epithelium [145].

Ca(<sup>2+</sup>)-ATPase activity was approximately 50% less in membranes prepared from cataractous lenses in comparison to clear lenses [146]. Marian et al. reported that the increased calcium concentration in cataractous lens was attributed to the increase of calcium-ATPase expression in its plasma membrane [147]. Activation of Ca-dependent enzymes may cause irreversible breakdown of structural proteins and apparent cell death of lens cells [148].

Excessive levels of Ca2+ in human lenses with cortical cataract have been found to play a major role in the opacification process. Ca2 attained 10,000 fold increase in human aqueous humour inwardly directed gradient across the plasma membrane and these are needed to maintain reduction. Both Na/Ca2 exchanger (NCX) and plasma membrane Ca2-ATPase (PMCA) actively remove Ca2 from the cells, whereas in endoplasmic reticulum, the Ca2-ATPase (SERCA) facilitated the storage of Ca++. Adenosine triphosphatase was found to be more active in lens epithelial cells compared with that of NCX in fiber cells. Activation of these channels may lead to a rapid increase in pathological calcium overload which facilitates the formation of cataract [149]. Plasma membrane calcium ATPase (PMCA) mRNA and protein levels attained a considerable higher level in cataractous lens which might be a compensatory mechanism to overcome increased intracellular calcium levels [147]. Chen et al. observed ATPbinding cassette protein (ABCA3) in human lens capsule, choroidretinal pigment epithelium and retinal pigment epithelial cells [150]. Mutation of ABCA3 protein was found to be associated with cataractmicrocornea syndrome.

Lenticular copper, lead, cadmium and calcium levels were

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markedly increased in cataractous human lenses especially in males. Cataract formation in diabetic patients possessed a higher concentration of copper which coincides with copper-containing superoxide dismutase and liberation of free radicals. Increased copper ions were attributed to its release from copper-containing enzymes affected by oxidative stress of hyperglycaemia and free radicals [151].

Maintenance of calcium homeostasis is important for the clarity of the lens.  $Ca(^{2+})$ -ATPase play a great role for the removal of cytosolic calcium, either across the plasma membrane or through intracellular organelles such as the endoplasmic reticulum. The activity of  $Ca(^{2+})$ -ATPase reached approximately 50% in membranes prepared from nuclear subcapsular, nuclear and brunescent cataracts [146].

The lenticular  $Ca^{2+}$  was found to be maintained by either plasma membrane  $Ca^{2+}$  pumps [152], or plasma membrane Na<sup>+</sup>: $Ca^{2+}$ exchangers [153], and endoplasmic reticulum  $Ca^{2+}$  pumps [152]. Cataractous lenses as a result of increased calcium concentration were observed in 14 out of 406 affected human lenses [154]. Odeigah and Patmore detected epithelial hyperplasia in cataractous lens of HY-1 genotype chick lenses and attributed it to the increased calcium binding capacity and elevated sialic acid [155]. Kuck and Kuck mentioned that mouse cataract was different from human senile cataracts due to invariable accumulation of calcium depending on the severity of cataract formation [156].

Cataractous lenses possessed abnormal intracellular ionic environment associated with increased levels of calcium (Ca<sup>2+</sup>), sodium (Na<sup>+</sup>), magnesium (Mg<sup>2+</sup>) and potassium (K<sup>+</sup>), Mg<sup>2+</sup> deficiency was found to enhances nitrous oxide production via inducible nitric oxide synthase in the lens and accelerate the progression of lens opacification [157,158]. Increased calcium ions in cataract [159], may activate calpain and transglutaminase, the markers of cataracts [160].

Lenticular cataracts were found to exhibit abnormal intracellular ions including increased level of calcium and sodium and depletion of magnesium and potassium. These were associated with a decrease of ATP and adenosine triphosphatase activity as well as increase of nitrous oxide production [158].

Depletion of Cu, Zn-SOD and catalase (CAT) activities associated with increase of malodialdhyde were detected in diabetic and senile cataractous patients [161].

Increase zinc, copper, and calcium; and a decrease in potassium concentration have a great role in cataractous lenses [162]. Copper ions attained a higher concentration in cataractous lenses of diabetic patients due to decrease in the reactivity of the copper-containing enzyme superoxide dismutase and an increase in hydrogen peroxide concentration [151].

# Antioxidant defense

Ocular tissues contain antioxidants that protect it from oxidative stress of free radicals such as redox antioxidant enzymes (catalase, superoxide dismutase, GSH peroxidase, glutathione S-transferase), ascorbic acid, glutathione, amino acids (cysteine and tyrosine) etc., [163,164] that scavenge free radicals. The antioxidant redox system was detected in the epithelial layer and outer cortical regions.

Fecondo and Augusteyn reported a marked depletion of

superoxide dismutase and glutathione peroxidase in the nuclear cataract of human, calf, rabbit and rat lenses [165]. Cataract formation is believed to be attributed to oxidative stress and reduction of the antioxidant defense, especially vitamin C concentration. Vitamin C was found to initiate glucose to protein glycation. Lens vitamin C concentration attained marked depletion in severe brown cataracts (88 mumol/100 g lens in mild cataracts, and 50 mumol/100 g in dark brown lenses) [109].

There was a detected reduction of the blood antioxidant redox chain and increase of thiobarbituric acid was found to be indicator of cataractous lenses in diabetic [166] and senile cataractous patients [167-169]. Jain and Bulakh confirmed similar findings of depletion in glutathione peroxidase and increase of thiobarbituric acid in cataractous lenses [170].

Lipid peroxidation (LPO) represent the main cause of cataract formation, initiated by enhanced production of oxygen free radicals in the eye fluids and tissues and inhibited antioxidant enzymatic and non-enzymatic defenses of the lens. The LPO products (diene conjugates, lipid hydroperoxides) and end products of fluorescent LPO were markedly detected in the aqueous humor samples of cataractous patients [171]. Lipid peroxidation led to degradation of the double bonds of unsaturated membrane hydrocarbon and altered the carbonyl and phosphate-oxygen sites of the fiber cell membranes [172]. These oxidative damages initiated in the absence of antioxidant defense [173].

The formation of cataract is believed to have resulted from an oxidative insult which decreases the antioxidant defense of the lens, especially vitamin C. The lens vitamin C concentration significantly decreased with cataract severity, but mostly in severe brown cataracts [109].

The authors finally concluded that both congenital and senile cataractous lenses resulted from multifactorial agents included genetic, biochemical transformations and alterations in antioxidant defences

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